RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: _10/540,394_

Source: _PCT_

Date Processed by STIC: _9/8/05_

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10 540 394   CRF Edit Date: 7/8/05
                                    Edited by: ___

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

✓ Deleted: ✓ Invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

____ Other:

_____
_____
_____

Revised 09/09/2003

PCT

RAW SEQUENCE LISTING   DATE: 07/08/2005
PATENT APPLICATION: US/10/540,394   TIME: 11:44:18

Input Set : **A:\pto.kd.txt**
Output Set: **N:\CRF4\07082005\J540394.raw**

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 3 <110> APPLICANT: SUNAHARA, Eiji
 4        ISHII, Takafumi
 5        YAMAMOTO, Koji
 6        SATO, Shuji
 8 <120> TITLE OF INVENTION: Novel protein and its use
10 <130> FILE REFERENCE: 3132US0P
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C--> `12 <140> CURRENT APPLICATION NUMBER: US/10/540,394`
C--> `12 <141> CURRENT FILING DATE: 2005-06-23`
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12 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/016655
13 <151> PRIOR FILING DATE: 2003-12-25
15 <150> PRIOR APPLICATION NUMBER: JP2002-378052
16 <151> PRIOR FILING DATE: 2002-12-26
18 <150> PRIOR APPLICATION NUMBER: JP2003-65497
19 <151> PRIOR FILING DATE: 2003-03-11
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24 <211> LENGTH: 837
25 <212> TYPE: PRT
26 <213> ORGANISM: Human
28 <400> SEQUENCE: 1
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32 Gly Ala Leu Pro Pro Arg Pro Pro Leu Leu Leu Leu Leu Leu Leu Leu
33                20                  25                  30
34 Leu Leu Leu Gln Pro Pro Pro Pro Thr Trp Ala Leu Ser Pro Arg Ile
35             35                  40                  45
36 Ser Leu Pro Leu Gly Ser Glu Glu Arg Pro Phe Leu Arg Phe Glu Ala
37         50                  55                  60
38 Glu His Ile Ser Asn Tyr Thr Ala Leu Leu Leu Ser Arg Asp Gly Arg
39 65                  70                  75                  80
40 Thr Leu Tyr Val Gly Ala Arg Glu Ala Leu Phe Ala Leu Ser Ser Asn
41                 85                  90                  95
42 Leu Ser Phe Leu Pro Gly Gly Glu Tyr Gln Glu Leu Leu Trp Gly Ala
43             100                 105                 110
44 Asp Ala Glu Lys Lys Gln Gln Cys Ser Phe Lys Gly Lys Asp Pro Gln
45         115                 120                 125
46 Arg Asp Cys Gln Asn Tyr Ile Lys Ile Leu Leu Pro Leu Ser Gly Ser
47     130                 135                 140
48 His Leu Phe Thr Cys Gly Thr Ala Ala Phe Ser Pro Met Cys Thr Tyr
49 145                 150                 155                 160
50 Ile Asn Met Glu Asn Phe Thr Leu Ala Arg Asp Glu Lys Gly Asn Val
51             165                 170                 175
52 Leu Leu Glu Asp Gly Lys Gly Arg Cys Pro Phe Asp Pro Asn Phe Lys
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Input Set : A:\pto.kd.txt
Output Set: N:\CRF4\07082005\J540394.raw

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55          195               200               205
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57       210               215               220
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59 225 .             230               235               240
60 Val Ala Ser Ala Tyr Ile Pro Glu Ser Leu Gly Ser Leu Gln Gly Asp
61                 245               250               255
62 Asp Asp Lys Ile Tyr Phe Phe Phe Ser Glu Thr Gly Gln Glu Phe Glu
63             260               265               270
64 Phe Phe Glu Asn Thr Ile Val Ser Arg Ile Ala Arg Ile Cys Lys Gly
65          275               280               285
66 Asp Glu Gly Gly Glu Arg Val Leu Gln Gln Arg Trp Thr Ser Phe Leu
67       290               295               300
68 Lys Ala Gln Leu Leu Cys Ser Arg Pro Asp Asp Gly Phe Pro Phe Asn
69 305             310               315               320
70 Val Leu Gln Asp Val Phe Thr Leu Ser Pro Ser Pro Gln Asp Trp Arg
71             325               330               335
72 Asp Thr Leu Phe Tyr Gly Val Phe Thr Ser Gln Trp His Arg Gly Thr
73          340               345               350
74 Thr Glu Gly Ser Ala Val Cys Val Phe Thr Met Lys Asp Val Gln Arg
75          355               360               365
76 Val Phe Ser Gly Leu Tyr Lys Glu Val Asn Arg Glu Thr Gln Gln Met
77       370               375               380
78 Val His Arg Asp Pro Pro Val Pro Thr Pro Arg Pro Gly Ala Cys Ile
79 385             390 .             395               400
80 Thr Asn Ser Ala Arg Glu Arg Lys Ile Asn Ser Ser Leu Gln Leu Pro
81 ·           405               410               415
82 Asp Arg Val Leu Asn Phe Leu Lys Asp His Phe Leu Met Asp Gly Gln
83          420               425               430
84 Val Arg Ser Arg Met Leu Leu Leu Gln Pro Gln Ala Arg Tyr Gln Arg
85          435               440               445
86 Val Ala Val His Arg Val Pro Gly Leu His His Thr Tyr Asp Val Leu
87       450               455               460
88 Phe Leu Gly Thr Gly Asp Gly Arg Leu His Lys Ala Val Ser Val Gly
89 465             470               475               480
90 Pro Arg Val His Ile Ile Glu Glu Leu Gln Ile Phe Ser Ser Gly Gln
91             485               490               495
92 Pro Val Gln Asn Leu Leu Leu Asp Thr His Arg Gly Leu Leu Tyr Ala
93          500               505               510
94 Ala Ser His Ser Gly Val Val Gln Val Pro Met Ala Asn Cys Ser Leu
95          515               520               525
96 Tyr Arg Ser Cys Gly Asp Cys Leu Leu Ala Arg Asp Pro Tyr Cys Ala
97       530               535               540
98 Trp Ser Gly Ser Ser Cys Lys His Val Ser Leu Tyr Gln Pro Gln Leu
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100 Ala Thr Arg Pro Trp Ile Gln Asp Ile Glu Gly Ala Ser Ala Lys Asp
101             565               570               575
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103             580             585             590
104 Glu Lys Pro Cys Glu Gln Val Gln Phe Gln Pro Asn Thr Val Asn Thr
105         595             600             605
106 Leu Ala Cys Pro Leu Leu Ser Asn Leu Ala Thr Arg Leu Trp Leu Arg
107     610             615             620
108 Asn Gly Ala Pro Val Asn Ala Ser Ala Ser Cys His Val Leu Pro Thr
109 625             630             635             640
110 Gly Asp Leu Leu Leu Val Gly Thr Gln Gln Leu Gly Glu Phe Gln Cys
111             645             650             655
112 Trp Ser Leu Glu Glu Gly Phe Gln Gln Leu Val Ala Ser Tyr Cys Pro
113             660             665             670
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115         675             680             685
116 Val Pro Val Ile Ile Ser Thr Ser Arg Val Ser Ala Pro Ala Gly Gly
117     690             695             700
118 Lys Ala Ser Trp Gly Ala Asp Arg Ser Tyr Trp Lys Glu Phe Leu Val
119 705             710             715             720
120 Met Cys Thr Leu Phe Val Leu Ala Val Leu Leu Pro Val Leu Phe Leu
121             725             730             735
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123             740             745             750
124 Cys Ala Ser Val His Pro Lys Thr Cys Pro Val Val Leu Pro Pro Glu
125         755             760             765
126 Thr Arg Pro Leu Asn Gly Leu Gly Pro Pro Ser Thr Pro Leu Asp His
127     770             775             780
128 Arg Gly Tyr Gln Ser Leu Ser Asp Ser Pro Pro Gly Ser Arg Val Phe
129 785             790             795             800
130 Thr Glu Ser Glu Lys Arg Pro Leu Ser Ile Gln Asp Ser Phe Val Glu
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135         835
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139 <212> TYPE: DNA
140 <213> ORGANISM: Human
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146 acctgggcgc tcagcccccg gatcagcctg cctctgggct ctgaagagcg gccattcctc     180
147 agattcgaag ctgaacacat ctccaactac acagcccttc tgctgagcag ggatggcagg     240
148 accctgtacg tgggtgctcg agaggccctc tttgcactca gtagcaacct cagcttcctg     300
149 ccaggcgggg agtaccagga gctgctttgg ggtgcagacg cagagaagaa acagcagtgc     360
150 agcttcaagg gcaaggaccc cagcgcgac tgtcaaaact acatcaagat cctcctgccg     420
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156 gtggcctcag cctacattcc tgagagcctg ggcagcttgc aaggcgatga tgacaagatc    780
157 tactttttct tcagcgagac tggccaggaa tttgagttct ttgagaacac cattgtgtcc    840
158 cgcattgccc gcatctgcaa gggcgatgag ggtggagagc gggtgctaca gcagcgctgg    900
159 acctccttcc tcaaggccca gctgctgtgc tcacggcccg acgatggctt ccccttcaac    960
160 gtgctgcagg atgtcttcac gctgagcccc agccccagg actggcgtga cacccttttc   1020
161 tatggggtct tcacttccca gtggcacagg ggaactacag aaggctctgc cgtctgtgtc   1080
162 ttcacaatga aggatgtgca gagagtcttc agcggcctct acaaggaggt gaaccgtgag   1140
163 acacagcaga tggtacaccg tgacccaccc gtgcccacac cccggcctgg agcgtgcatc   1200
164 accaacagtg cccgggaaag gaagatcaac tcatccctgc agctcccaga ccgcgtgctg   1260
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166 cagccccagg ctcgctacca gcgcgtggct gtacaccgcg tccctggcct gcaccacacc   1380
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177 gagggcttcc agcagctggt agccagctac tgcccagagg tggtggagga cggggtggca   2040
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199 cggccaccgc tgctgctgct cctgctgctg ctgctcctgc tgcagccgcc gcctccgacc    360
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204 ttcaagggca aggacccaca gcgcgactgt caaaactaca tcaagatcct cctgccgctc    660
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VERIFICATION SUMMARY   DATE: 07/08/2005
PATENT APPLICATION: **US/10/540,394**   TIME: 11:44:19

Input Set : **A:\pto.kd.txt**
Output Set: **N:\CRF4\07082005\J540394.raw**

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

Raw Sequence Listing before editing,
for reference only

PCT

RAW SEQUENCE LISTING   DATE: 07/01/2005
PATENT APPLICATION: US/10/540,394   TIME: 14:40:43

        Input Set : A:\3132-SEQ.txt
        Output Set: N:\CRF4\07012005\J540394.raw

     3 <110> APPLICANT: SUNAHARA, Eiji
     4   ISHII, Takafumi
     5   YAMAMOTO, Koji
     6   SATO, Shuji
     8 <120> TITLE OF INVENTION: Novel protein and its use
    10 <130> FILE REFERENCE: 3132USOP
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/540,394
C--> 12 <141> CURRENT FILING DATE: 2005-06-23
    12 <150> PRIOR APPLICATION NUMBER: PCT/JP2003/016655
    13 <151> PRIOR FILING DATE: 2003-12-25
    15 <150> PRIOR APPLICATION NUMBER: JP2002-378052
    16 <151> PRIOR FILING DATE: 2002-12-26
    18 <150> PRIOR APPLICATION NUMBER: JP2003-65497
    19 <151> PRIOR FILING DATE: 2003-03-11
    21 <160> NUMBER OF SEQ ID NOS: 25

Does Not Comply
Corrected Diskette Needed
(pg. 1)

ERRORED SEQUENCES

    1101 <210> SEQ ID NO: 25
    1102 <211> LENGTH: 14
    1103 <212> TYPE: PRT
    1104 <213> ORGANISM: Human
    1106 <400> SEQUENCE: 25
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    1109   5   10
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VERIFICATION SUMMARY   DATE: 07/01/2005
PATENT APPLICATION: **US/10/540,394**   TIME: 14:40:44

Input Set : **A:\3132-SEQ.txt**
Output Set: **N:\CRF4\07012005\J540394.raw**

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1113 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:25